

D. Clark

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Page 1 of 6
NOV 15 2000
1633
TECH CENTER 1600/2300

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/245,603A
DATE: 11/03/2000
TIME: 15:41:24

Input Set : A:\PTO.txt
Output Set: N:\CRF3\11032000\I245603A.raw

3 <110> APPLICANT: Curiel, David T.
4 Krasnykh, Victor N.
5 Dmitriev, Igor
7 <120> TITLE OF INVENTION: Adenovirus Vector Containing A Heterologous Peptide
8 Epitope in the HI Loop of the Fiber Knob
W--> 9 <130> FILE REFERENCE: D6080
W--> 10 <140> CURRENT APPLICATION NUMBER: 09/245,603A
11 <141> CURRENT FILING DATE: 1999-02-05
12 <150> PRIOR APPLICATION NUMBER: US 60/099,801
14 <151> PRIOR FILING DATE: 1998-09-10
W--> 16 <160> NUMBER OF SEQ ID: 17
18 <210> SEQ ID NO: 1
19 <211> LENGTH: 38
20 <212> TYPE: DNA
21 <213> ORGANISM: artificial sequence
W--> 22 <220> FEATURE:
23 <221> NAME/KEY: primer_bind
24 <223> OTHER INFORMATION: Forward primer F1 used to generate a gene encoding
25 the Ad5 fiber knob domain with the HI loop deleted.
W--> 26 <400> SEQUENCE: 1
27 taaggatccg gtgccattac agtaggaac aaaaataa 38
29 <210> SEQ ID NO: 2
30 <211> LENGTH: 43
31 <212> TYPE: DNA
32 <213> ORGANISM: artificial sequence
W--> 33 <220> FEATURE:
34 <221> NAME/KEY: primer_bind
35 <223> OTHER INFORMATION: Reverse primer R1 used to generate a gene encoding
36 the Ad5 fiber knob domain with the HI loop deleted.
W--> 37 <400> SEQUENCE: 2
38 catagagtat gcagatatcg ttagtggttac aggttttagtt ttg 43
40 <210> SEQ ID NO: 3
41 <211> LENGTH: 42
42 <212> TYPE: DNA
43 <213> ORGANISM: artificial sequence
W--> 44 <220> FEATURE:
45 <221> NAME/KEY: primer_bind
46 <223> OTHER INFORMATION: Forward primer F2 used to generate a gene encoding
47 the Ad5 fiber knob domain with the HI loop deleted.
W--> 48 <400> SEQUENCE: 3
49 gtaacactaa cgatatctgc atactctatg tcattttcat gg 42
51 <210> SEQ ID NO: 4
52 <211> LENGTH: 41
53 <212> TYPE: DNA
54 <213> ORGANISM: artificial sequence
W--> 55 <220> FEATURE:
56 <221> NAME/KEY: primer_bind

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57 <223> OTHER INFORMATION: Reverse primer R2 used to generate a gene encoding
58   the Ad5 fiber knob domain with the HI loop deleted.
W--> 59 <400> SEQUENCE: 4
60   cccaagctta caattgaaaa ataaacacgt tgaaacataa c      41
62 <210> SEQ ID NO: 5
63 <211> LENGTH: 63
64 <212> TYPE: DNA
65 <213> ORGANISM: artificial sequence
W--> 66 <220> FEATURE:
67 <223> OTHER INFORMATION: Oligonucleotide annealed with SEQ ID NO: 6 to form a
68   duplex and cloned into EcoRV-digested pQE.KNOBDHI.
W--> 69 <400> SEQUENCE: 5
70   tacactaaac ggtacccagg aaacaggaga cacaactgac tacaaggacg acgatgacaa 60
71   gcc                                           63
73 <210> SEQ ID NO: 6
74 <211> LENGTH: 63
75 <212> TYPE: DNA
76 <213> ORGANISM: artificial sequence
W--> 77 <220> FEATURE:
78 <223> OTHER INFORMATION: Oligonucleotide annealed with SEQ ID NO: 5 to form a
79   duplex and cloned into EcoRV-digested pQE.KNOBDHI.
W--> 80 <400> SEQUENCE: 6
81   ggcttgatc cgtcgtcctt gtagtcagtt gtgtctctctg tttctctgggt accggttagt 60
82   gta                                           63
84 <210> SEQ ID NO: 7
85 <211> LENGTH: 29
86 <212> TYPE: DNA
87 <213> ORGANISM: artificial sequence
W--> 88 <220> FEATURE:
89 <223> OTHER INFORMATION: Oligonucleotide used in synthetic duplex which
90   encodes MetHis6Lys.
W--> 91 <400> SEQUENCE: 7
92   gatccatgca tcaccatcac catcacaag      29
94 <210> SEQ ID NO: 8
95 <211> LENGTH: 29
96 <212> TYPE: DNA
97 <213> ORGANISM: artificial sequence
W--> 98 <220> FEATURE:
99 <223> OTHER INFORMATION: Oligonucleotide used in synthetic duplex which
100   encodes MetHis6Lys.
W--> 101 <400> SEQUENCE: 8
102  cgcgcttgatg atggtgatgg tgatgcatg      29
104 <210> SEQ ID NO: 9
105 <211> LENGTH: 16
106 <212> TYPE: DNA
107 <213> ORGANISM: artificial sequence
W--> 108 <220> FEATURE:
109 <223> OTHER INFORMATION: An NdeI-SwaI linker ligated to plasmid pTG3602 after
110   partial digestion of the plasmid with NdeI.

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W--> 111 <400> SEQUENCE: 9
112 taccatttta aatggg          16
114 <210> SEQ ID NO: 10
115 <211> LENGTH: 66
116 <212> TYPE: DNA
117 <213> ORGANISM: artificial sequence
W--> 118 <220> FEATURE:
119 <223> OTHER INFORMATION: Oligonucleotide in duplex cloned into EcoRV site
120   of plasmid pQE.KNOBDHI generating pQE.KNOB.RGDHI.
W--> 121 <400> SEQUENCE: 10
122 cacactaaac ggtacacagg aaacagygaga cacaacttgt gactgcccgc gagactgttt 60
123 ctgccc                      66
125 <210> SEQ ID NO: 11
126 <211> LENGTH: 66
127 <212> TYPE: DNA
128 <213> ORGANISM: artificial sequence
W--> 129 <220> FEATURE:
130 <221> NAME/KEY: primer_bind
131 <223> OTHER INFORMATION: Oligonucleotide in duplex cloned into EcoRV site
132   of plasmid pQE.KNOBDHI generating pQE.KNOB.RGDHI.
W--> 133 <400> SEQUENCE: 11
134 gggcagaaac agtctccgcg gcagtcacaa gttgtgtctc ctgtttcctg tgtaccgttt 60
135 agtgtg                      66
137 <210> SEQ ID NO: 12
138 <211> LENGTH: 41
139 <212> TYPE: DNA
140 <213> ORGANISM: artificial sequence
W--> 141 <220> FEATURE:
142 <223> OTHER INFORMATION: Oligonucleotide in synthetic duplex used to
143   replace 41 bp PacI-ClaI-fragment in pcDNA.Luc,
144   generating pcLucPCL.
W--> 145 <400> SEQUENCE: 12
146 caaatataaa ggatatcagg tggccccgcg tgaattggag t          41
148 <210> SEQ ID NO: 13
149 <211> LENGTH: 45
150 <212> TYPE: DNA
151 <213> ORGANISM: artificial sequence
W--> 152 <220> FEATURE:
153 <223> OTHER INFORMATION: Oligonucleotide in synthetic duplex used to
154   replace 41 bp PacI-ClaI-fragment in pcDNA.Luc,
155   generating pcLucPCL.
W--> 156 <400> SEQUENCE: 13
157 cgactccaat tcagcggggg ccacctgata tcctttgtat ttgat          45
159 <210> SEQ ID NO: 14
160 <211> LENGTH: 13
161 <212> TYPE: PRT
162 <213> ORGANISM: artificial sequence
W--> 163 <220> FEATURE:
164 <223> OTHER INFORMATION: Amino acid sequence deleted from the HI loop of

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165         the fiber knob domain and replaced with a
166         unique EcoRV site.
W--> 167 <400> SEQUENCE: 14
168 Thr Leu Asn Gly Thr Gln Glu Thr Gly Asp Thr Thr Pro
169             5             10
171 <210> SEQ ID NO: 15
172 <211> LENGTH: 8
173 <212> TYPE: PRT
174 <213> ORGANISM: artificial sequence
W--> 175 <220> FEATURE:
176 <223> OTHER INFORMATION: Amino acid sequence of the FLAG octapeptide.
W--> 177 <400> SEQUENCE: 15
178 Asp Tyr Lys Asp Asp Asp Asp Lys
179             5
181 <210> SEQ ID NO: 16
182 <211> LENGTH: 9
183 <212> TYPE: PRT
184 <213> ORGANISM: artificial sequence
W--> 185 <220> FEATURE:
186 <223> OTHER INFORMATION: Amino acid sequence of a RGD peptide incorporated
187         into the region of the fiber gene within the HI loop.
W--> 188 <400> SEQUENCE: 16
189 Cys Asp Cys Arg Gly Asp Cys Phe Cys
190             5
192 <210> SEQ ID NO: 17
193 <211> LENGTH: 13
194 <212> TYPE: PRT
195 <213> ORGANISM: artificial sequence
W--> 196 <220> FEATURE:
197 <223> OTHER INFORMATION: Amino acid sequence of peptide replacing the
198         RGD coding sequence.
W--> 199 <400> SEQUENCE: 17
200 Thr Leu Asn Gly Thr Gln Glu Thr Gly Asp Thr Thr Pro
201             5             10

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VERIFICATION SUMMARY

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L:10 M:283 W: Missing Blank Line separator, <140> field identifier
L:16 M:283 W: Missing Blank Line separator, <160> field identifier
L:22 M:283 W: Missing Blank Line separator, <220> field identifier
L:26 M:283 W: Missing Blank Line separator, <400> field identifier
L:33 M:283 W: Missing Blank Line separator, <220> field identifier
L:37 M:283 W: Missing Blank Line separator, <400> field identifier
L:44 M:283 W: Missing Blank Line separator, <220> field identifier
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